RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/577, 390_

Source: _JFWO_

Date Processed by STIC: _05/11/2006_

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: _10/577, 390_   CRF Edit Date: _05/11/2006_
                                    Edited by: ___DH___

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

__/__ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

____ Other:

_____
_____
_____

Revised 09/09/2003

IFW ℘

RAW SEQUENCE LISTING
PATENT APPLICATION: **US/10/577,390**

DATE: 05/11/2006
TIME: 13:13:59

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\05092006\J577390.raw**

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 4 <110> APPLICANT: Chen, Bao-Lu
 5        Hurst, Deborah
 6        Lee, Sang Hoon
 7        Long, Li
 8        Lu, Xiaofeng
 9        Luqman, Mohammad
10        Yabannavar, Asha
11        Zaror, Isabel
13 <120> TITLE OF INVENTION: Antagonist Anti-CD40 Monoclonal
14        Antibodies and Methods for Their Use
17 <130> FILE REFERENCE: PP20107.004 (282916)
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C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/577,390
C--> 19 <141> CURRENT FILING DATE: 2006-04-27
     19 <150> PRIOR APPLICATION NUMBER: 60/565,710
     20 <151> PRIOR FILING DATE: 2004-04-27
     22 <150> PRIOR APPLICATION NUMBER: 60/525,579
     23 <151> PRIOR FILING DATE: 2003-11-26
     25 <150> PRIOR APPLICATION NUMBER: 60/517,337
     26 <151> PRIOR FILING DATE: 2003-11-04
     28 <160> NUMBER OF SEQ ID NOS: 12
     30 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     32 <210> SEQ ID NO: 1
     33 <211> LENGTH: 720
     34 <212> TYPE: DNA
     35 <213> ORGANISM: Artificial Sequence
     37 <220> FEATURE:
     38 <223> OTHER INFORMATION: Coding sequence for light chain of 12.12 human
     39        anti-CD40 antibody
W--> 41 <221> NAME/KEY: CDS
     42 <222> LOCATION: (1)...(720)
W--> 44 <400> 1
     45 atg gcg ctc cct gct cag ctc ctg ggg ctg cta atg ctc tgg gtc tct    48
     46 Met Ala Leu Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Ser
     47  1               5                  10                  15
     49 gga tcc agt ggg gat att gtg atg act cag tct cca ctc tcc ctg acc    96
     50 Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Thr
     51            20                  25                  30
     53 gtc acc cct gga gag ccg gcc tcc atc tcc tgc agg tcc agt cag agc   144
     54 Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
     55          35                  40                  45
     57 ctc ctg tat agt aat gga tac aac tat ttg gat tgg tac ctg cag aag   192
     58 Leu Leu Tyr Ser Asn Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys
     59       50                  55                  60
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RAW SEQUENCE LISTING   DATE: 05/11/2006
PATENT APPLICATION: US/10/577,390   TIME: 13:13:59

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\05092006\J577390.raw

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61 cca ggg cag tct cca cag gtc ctg atc tct ttg ggt tct aat cgg gcc    240
62 Pro Gly Gln Ser Pro Gln Val Leu Ile Ser Leu Gly Ser Asn Arg Ala
63    65              70               75                  80
65 tcc ggg gtc cct gac agg ttc agt ggc agt gga tca ggc aca gat ttt    288
66 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
67            85               90                  95
69 aca ctg aaa atc agc aga gtg gag gct gag gat gtt ggg gtt tat tac    336
70 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
71        100              105                 110
73 tgc atg caa gct cga caa act cca ttc act ttc ggc cct ggg acc aaa    384
74 Cys Met Gln Ala Arg Gln Thr Pro Phe Thr Phe Gly Pro Gly Thr Lys
75        115              120                 125
77 gtg gat atc aga cga act gtg gct gca cca tct gtc ttc atc ttc ccg    432
78 Val Asp Ile Arg Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro
79    130              135                 140
81 cca tct gat gag cag ttg aaa tct gga act gcc tct gtt gtg tgc ctg    480
82 Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu
83 145              150                 155                 160
85 ctg aat aac ttc tat ccc aga gag gcc aaa gta cag tgg aag gtg gat    528
86 Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp
87            165              170                 175
89 aac gcc ctc caa tcg ggt aac tcc cag gag agt gtc aca gag cag gac    576
90 Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp
91        180              185                 190
93 agc aag gac agc acc tac agc ctc agc agc acc ctg acg ctg agc aaa    624
94 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys
95        195              200                 205
97 gca gac tac gag aaa cac aaa gtc tac gcc tgc gaa gtc acc cat cag    672
98 Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln
99    210              215                 220
101 ggc ctg agc tcg ccc gtc aca aag agc ttc aac agg gga gag tgt tag    720
102 Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys  *
103 225              230                 235
107 <210> SEQ ID NO: 2
108 <211> LENGTH: 239
109 <212> TYPE: PRT
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Light chain of 12.12 human anti-CD40 antibody
115 <400> SEQUENCE: 2
116 Met Ala Leu Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Ser
117  1              5                10                  15
118 Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Thr
119            20               25                  30
120 Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
121        35               40                  45
122 Leu Leu Tyr Ser Asn Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys
123    50               55                  60
124 Pro Gly Gln Ser Pro Gln Val Leu Ile Ser Leu Gly Ser Asn Arg Ala
```

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125 65                     70                75                80
126 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
127                    85                90                95
128 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
129                    100               105               110
130 Cys Met Gln Ala Arg Gln Thr Pro Phe Thr Phe Gly Pro Gly Thr Lys
131.                   115               120               125
132 Val Asp Ile Arg Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro
133     130               135               140
134 Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu
135 145               150               155               160
136 Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp
137                    165               170               175
138 Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp
139                    180               185               190
140 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys
141     195               200               205
142 Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln
143.    210               215               220
144 Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
145 225               230               235
148 <210> SEQ ID NO: 3
149 <211> LENGTH: 2016
150 <212> TYPE: DNA
151 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: Coding sequence for heavy chain of 12.12 human
155       anti-CD40 antibody (with introns)
157 <400> SEQUENCE: 3
158 atg gag ttt ggg ctg agc tgg gtt ttc ctt gtt gct att tta aga ggt    48
159 gtc cag tgt cag gtg cag ttg gtg gag tct ggg gga ggc gtg gtc cag    96
160 cct ggg agg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc   144
161 agt agc tat ggc atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg   192
162 gag tgg gtg gca gtt ata tca tat gag gaa agt aat aga tac cat gca   240
163 gac tcc gtg aag ggc cga ttc acc atc tcc aga gac aat tcc aag atc   288
164 acg ctg tat ctg caa atg aac agc ctc aga act gag gac acg gct gtg   336
165 tat tac tgt gcg aga gat ggg ggt ata gca gca cct ggg cct gac tac   384
166 tgg ggc cag gga acc ctg gtc acc gtc tcc tca gca agt acc aag ggc   432
167 cca tcc gtc ttc ccc ctg gcg ccc gct agc aag agc acc tct ggg ggc   480
168 aca gcg gcc ctg ggc tgc ctg gtc aag gac tac ttc ccc gaa ccg gtg   528
169 acg gtg tcg tgg aac tca ggc gcc ctg acc agc ggc gtg cac acc ttc   576
170 ccg gct gtc cta cag tcc tca gga ctc tac tcc ctc agc agc gtg gtg   624
171 acc gtg ccc tcc agc agc ttg ggc acc cag acc tac atc tgc aac gtg   672
172 aat cac aag ccc agc aac acc aag gtg gac aag aga gtt ggt gag agg   720
173 cca gca cag gga ggg agg gtg tct gct gga agc cag gct cag cgc tcc   768
174 tgc ctg gac gca tcc cgg cta tgc agt ccc agt cca ggg cag caa ggc   816
175 agg ccc cgt ctg cct ctt cac ccg gag gcc tct gcc cgc ccc act cat   864
176 gct cag gga gag ggt ctt ctg gct ttt tcc cca ggc tct ggg cag gca   912
177 cag gct agg tgc ccc taa ccc agg ccc tgc aca caa agg ggc agg tgc   960
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178 tgg gct cag acc tgc caa gag cca tat ccg gga gga ccc tgc ccc tga   1008
179 cct aag ccc acc cca aag gcc aaa ctc tcc act ccc tca gct cgg aca   1056
180 cct tct ctc ctc cca gat tcc agt aac tcc caa tct tct ctc tgc aga   1104
181 gcc caa atc ttg tga caa aac tca cac atg ccc acc gtg ccc agg taa   1152
182 gcc agc cca ggc ctc gcc ctc cag ctc aag gcg gga cag gtg ccc tag   1200
183 agt agc ctg cat cca ggg aca ggc ccc agc cgg gtg ctg aca cgt cca   1248
184 cct cca tct ctt cct cag cac ctg aac tcc tgg ggg gac cgt cag tct   1296
185 tcc tct tcc ccc caa aac cca agg aca ccc tca tga tct ccc gga ccc   1344
186 ctg agg tca cat gcg tgg tgg tgg acg tga gcc acg aag acc ctg agg   1392
187 tca agt tca act ggt acg tgg acg gcg tgg agg tgc ata atg cca aga   1440
188 caa agc cgc ggg agg agc agt aca aca gca cgt acc gtg tgg tca gcg   1488
189 tcc tca ccg tcc tgc acc agg act ggc tga atg gca agg agt aca agt   1536
190 gca agg tct cca aca aag ccc tcc cag ccc cca tcg aga aaa cca tct   1584
191 cca aag cca aag gtg gga ccc gtg ggg tgc gag ggc cac atg gac aga   1632
192 ggc cgg ctc ggc cca ccc tct gcc ctg aga gtg acc gct gta cca acc   1680
193 tct gtc cct aca ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc   1728
194 cca tcc cgg gag gag atg acc aag aac cag gtc agc ctg acc tgc ctg   1776
195 gtc aaa ggc ttc tat ccc agc gac atc gcc gtg gag tgg gag agc aat   1824
196 ggg cag ccg gag aac aac tac aag acc acg cct ccc gtg ctg gac tcc   1872
197 gac ggc tcc ttc ttc ctc tat agc aag ctc acc gtg gac aag agc agg   1920
198 tgg cag cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg   1968
199 cac aac cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa tga   2016
202 <210> SEQ ID NO: 4
203 <211> LENGTH: 469
204 <212> TYPE: PRT
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Heavy chain of 12.12 human anti-CD40 antibody
210 <400> SEQUENCE: 4
211 Met Glu Phe Gly Leu Ser Trp Val Phe Leu Val Ala Ile Leu Arg Gly
212 1               5                   10                  15
213 Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
214             20                  25                  30
215 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
216         35                  40                  45
217 Ser Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
218     50                  55                  60
219 Glu Trp Val Ala Val Ile Ser Tyr Glu Glu Ser Asn Arg Tyr His Ala
220 65                  70                  75                  80
221 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Ile
222                 85                  90                  95
223 Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Thr Glu Asp Thr Ala Val
224             100                 105                 110
225 Tyr Tyr Cys Ala Arg Asp Gly Gly Ile Ala Ala Pro Gly Pro Asp Tyr
226         115                 120                 125
227 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly
228     130                 135                 140
229 Pro Ser Val Phe Pro Leu Ala Pro Ala Ser Lys Ser Thr Ser Gly Gly
230 145                 150                 155                 160
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231 Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val
232               165                 170                 175
233 Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe
234               180                 185                 190
235 Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val
236           195                 200                 205
237 Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val
238       210                 215                 220
239 Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Pro Lys
240 225                 230                 235                 240
241 Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu
242               245                 250                 255
243 Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
244           260                 265                 270
245 Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
246       275                 280                 285
247 Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val
248     290                 295                 300
249 Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
250 305                 310                 315                 320
251 Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
252               325                 330                 335
253 Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
254               340                 345                 350
255 Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
256       355                 360                 365
257 Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
258     370                 375                 380
259 Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
260 385                 390                 395                 400
261 Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
262               405                 410                 415
263 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
264           420                 425                 430
265 Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
266           435                 440                 445
267 Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
268     450                 455                 460
269 Leu Ser Pro Gly Lys
270 465
273 <210> SEQ ID NO: 5
274 <211> LENGTH: 469
275 <212> TYPE: PRT
276 <213> ORGANISM: Artificial Sequence
278 <220> FEATURE:
279 <223> OTHER INFORMATION: Heavy chain of variant of 12.12 human anti-CD40
280       antibody
282 <400> SEQUENCE: 5
283 Met Glu Phe Gly Leu Ser Trp Val Phe Leu Val Ala Ile Leu Arg Gly
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VERIFICATION SUMMARY　　　　　　　　DATE: 05/11/2006
PATENT APPLICATION: **US/10/577,390**　　TIME: 13:14:00

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\05092006\J577390.raw**

L:19 M:270 C: Current Application Number differs, Replaced Current Application No
L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:41 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:44 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:542 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:643 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11

Raw Sequence Listing before editing,
for reference only

IFWP

RAW SEQUENCE LISTING   DATE: 05/09/2006
PATENT APPLICATION: US/10/577,390   TIME: 11:14:57

           Input Set : A:\seqlst.txt
           Output Set: N:\CRF4\05092006\J577390.raw

   4 <110> APPLICANT: Chen, Bao-Lu
   5   Hurst, Deborah
   6   Lee, Sang Hoon
   7   Long, Li
   8   Lu, Xiaofeng
   9   Luqman, Mohammad
  10   Yabannavar, Asha
  11   Zaror, Isabel
  13 <120> TITLE OF INVENTION: Antagonist Anti-CD40 Monoclonal
  14   Antibodies and Methods for Their Use
  17 <130> FILE REFERENCE: PP20107.004 (282916)
C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/577,390
C--> 19 <141> CURRENT FILING DATE: 2006-04-27
  19 <150> PRIOR APPLICATION NUMBER: 60/565,710
  20 <151> PRIOR FILING DATE: 2004-04-27
  22 <150> PRIOR APPLICATION NUMBER: 60/525,579
  23 <151> PRIOR FILING DATE: 2003-11-26
  25 <150> PRIOR APPLICATION NUMBER: 60/517,337
  26 <151> PRIOR FILING DATE: 2003-11-04
  28 <160> NUMBER OF SEQ ID NOS: 12
  30 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply Corrected Diskette Needed (pg-2)

ERRORED SEQUENCES

 717 <210> SEQ ID NO: 12
 718 <211> LENGTH: 277
 719 <212> TYPE: PRT
 720 <213> ORGANISM: Homo sapiens
 722 <400> SEQUENCE: 12
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 724 1   5   10   15
 725 Ala Val His Pro Glu Pro Pro Thr Ala Cys Arg Glu Lys Gln Tyr Leu
 726   20   25   30
 727 Ile Asn Ser Gln Cys Cys Ser Leu Cys Gln Pro Gly Gln Lys Leu Val
 728   35   40   45
 729 Ser Asp Cys Thr Glu Phe Thr Glu Thr Glu Cys Leu Pro Cys Gly Glu
 730   50   55   60
 731 Ser Glu Phe Leu Asp Thr Trp Asn Arg Glu Thr His Cys His Gln His
 732 65   70   75   80
 733 Lys Tyr Cys Asp Pro Asn Leu Gly Leu Arg Val Gln Gln Lys Gly Thr
 734   85   90   95
 735 Ser Glu Thr Asp Thr Ile Cys Thr Cys Glu Glu Gly Trp His Cys Thr

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736             100             105          110
737 Ser Glu Ala Cys Glu Ser Cys Val Leu His Arg Ser Cys Ser Pro Gly
738         115             120             125
739 Phe Gly Val Lys Gln Ile Ala Thr Gly Val Ser Asp Thr Ile Cys Glu
740     130                 135             140
741 Pro Cys Pro Val Gly Phe Phe Ser Asn Val Ser Ser Ala Phe Glu Lys
742 145                 150             155             160
743 Cys His Pro Trp Thr Ser Cys Glu Thr Lys Asp Leu Val Val Gln Gln
744             165                 170             175
745 Ala Gly Thr Asn Lys Thr Asp Val Val Cys Gly Pro Gln Asp Arg Leu
746         180             185             190
747 Arg Ala Leu Val Val Ile Pro Ile Ile Phe Gly Ile Leu Phe Ala Ile
748         195             200             205
749 Leu Leu Val Leu Val Phe Ile Lys Lys Val Ala Lys Lys Pro Thr Asn
750     210             215             220
751 Lys Ala Pro His Pro Lys Gln Glu Pro Gln Glu Ile Asn Phe Pro Asp
752 225             230             235             240
753 Asp Leu Pro Gly Ser Asn Thr Ala Ala Pro Val Gln Glu Thr Leu His
754             245             250             255
755 Gly Cys Gln Pro Val Thr Gln Glu Asp Gly Lys Glu Ser Arg Ile Ser
756             260             265             270
757 Val Gln Glu Arg Gln
758             275
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E--> 760 12
     766 RTA01/2168223v1

deleted

VERIFICATION SUMMARY   DATE: 05/09/2006
PATENT APPLICATION: **US/10/577,390**   TIME: 11:14:58

Input Set : **A:\seqlst.txt**
Output Set: **N:\CRF4\05092006\J577390.raw**

L:19 M:270 C: Current Application Number differs, Replaced Current Application No
L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:41 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:44 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:542 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:643 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:760 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12